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Figure 1

TROM-DAJA mFas-IC 1225A GST-Binding to GST fusion proteins **TNCIFERASE** GST-mFas-IC GST-huFas-IC TAOM-DAJA TUCIFERASE

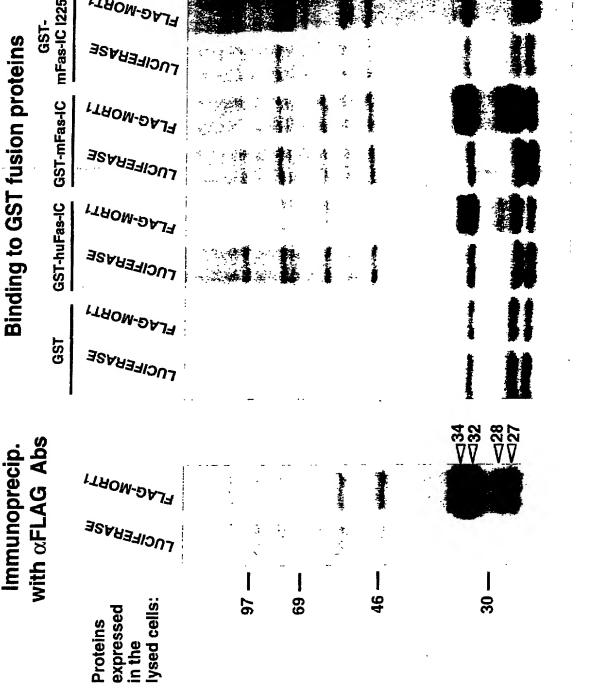
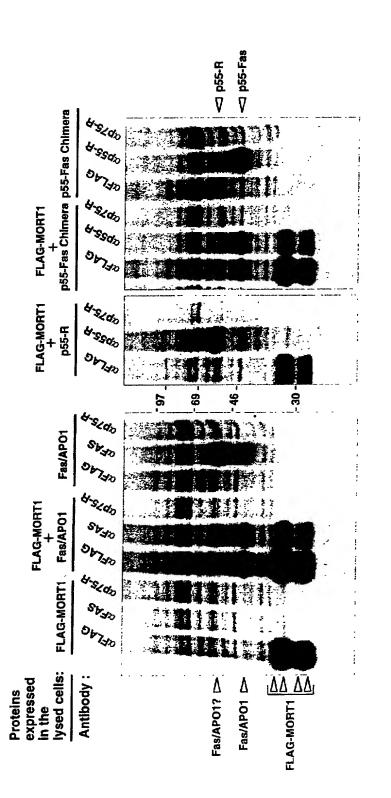


Figure 2



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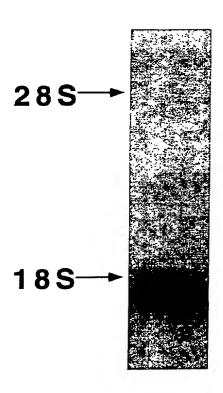


Figure 3

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1/1 GTG AAT CAG GCA CCG GAG TGC AGG TTC GGG GGT GGA ATC CTT GGG CCG CTG GGC AAG CGG V N Q A P E C R F G G G I L G P L G K R 61/21 91/31 61/21 CGA GAC CTG GCC AGG GCC AGG GAG CCG AGG ACA GAG GGC GCG CGG AGG GCC GGG CCG CAG R D L A R A S E P R T E G A R R A
121/41 151/51 G P CCC CGG CCG CTT GCA GAC CCC GCC ATG GAC CCG TTC CTG GTG CTG CAC TCG GTG TCG R P L A D P A M D P F L V L 211/71 181/61 TCC AGC CTG TCG AGC AGC GAG CTG ACC GAG CTC AAG TTC CTA TGC CTC GGG CGC GTG GTC S S L S S S E L T Ê L K F L С 271/91 AAG CCC AAG CTG GAG CGC GTG CAG AGC GGC CTA GAC CTC TTC TCC ATG CTG CTG GAG CAG RKLERVQSGLDLFSMLLEQ ./101 331/111 301/101 AAC GAC CTG GAG CCC GGG CAC ACC GAG CTC CTG CGC GAG CTG CTC GCC TCC CTG CGG CGC N D L E P G H T E L L
361/121 L R E L L A S 391/131 CAC GAC CTG CTG CGG CGC GTC GAC GAC TTC GAG GCG GGG GCG GCG GCG GCG GCG CCT L L R R V D D F E A G A A A G A A 451/151 421/141 GGG GAA GAA GAC CTG TGT GCA GCA TTT AAC GTC ATA TGT GAT AAT GTG GGG AAA GAT TGG E E D L C A A F N V C D N V G K D 511/171 481/161 AGA AGG CTG GCT CGT CAG CTC AAA GTC TCA GAC ACC AAG ATC GAC AGC ATC GAG GAC AGA R R L A R Q L K V S D T 541/181 571/191 T K I D S I E D R Y P R N L T E R 501/201 TAC CCC CGC AAC CTG ACA GAG CGT GTG CGG GAG TCA CTG AGA ATC TGG AAG AAC ACA GAG V R E S L R I W X N T 631/211 ANG GAG AND GCA ACA GTG GCC CAC CTG GTG GGG GCT CTC AGG TCC TGC CAG ATG AND CTG N A T V A H L V G G A L R S C Q M N L 691/231 661/221 GTG GCT GAC CTG GTA CAA GAG GTT CAG CAG GCC CGT GAC CTC CAG AAC AGG AGT GGG GCC <u>V</u> A D L V Q E V Q Q A R D L Q N R S G 721/241 751/251 ATG TCC CCG ATG TCA TGG AAC TCA GAC GCA TCT ACC TCC GAA GCG TCC TGA TGG GCC GCT S P M S W N S D A S S E 811/271 781/261 GCT TTG CGC TGG TGG ACC ACA GGC ATC TAC ACA GCC TGG ACT TTG GTT CTC TCC AGG AAG 841/281 871/291 GTA GCC CAG CAC TGT GAA GAC CCA GCA GGA AGC CAG GCT GAG TGA GCC ACA GAC CAC CTG 931/311 901/301 CTT CTG ANC TCA AGC TGC GTT TAT TAN TGC CTC TCC CGC ACC AGG CCG GGC TTG GGC CCT 991/331 961/321 GCA CAG ATA TTT CCA TTT CTT CCT CAC TAT GAC ACT GAG CAA GAT CTT GTC TCC ACT AAA 1051/351 1021/341 TGA GCT CCT GCG GGA GTA GTT GGA AAG TTG GAA CCG TGT CCA GCA CAG AAG GAA TCT GTG 1111/371 CAG ATG AGG AGT CAC ACT GTT ACT CCA CAG CGG AGG AGA CCA GCT CAG AGG CCC AGG AAT 1141/381 1171/391 CGG AGC GAA GCA GAG AGG TGG AGA ACT GGG ATT TGA ACC CCC GCC ATC CTT CAC CAG AGC 1201/401 1231/411 CCA TGC TCA ACC ACT GTG GCG TTC TGC TGC CCC TGC AGT TGG CAG AAA GGA TGT TTT TGT 1291/431 1261/421 CCC ATT TCC TTG GAG GCC ACC GGG ACA GAC CTG GAC ACT AGG GTC AGG CGG GGT GCT GTG 1351/451 1321/441 GTG GGG AGA GGC ATG GCT GGG GTG GGG GTG GGG AGA CCT GGT TGG CCG TGG TCC AGC TCT 1411/471 1381/461 TGG CCC CTG TGT GAG TTG AGT CTC CTC TCT GAG ACT GCT AAG TAG GGG CAG TGA TGG TTG 1441/481 1471/491 CCA GGA CGA ATT GAG ATA ATA TCT GTG AGG TGC TGA TGA GTG ATT GAC ACA CAG CAC TCT 1531/511 1501/501 CTA AAT CTT CCT TGT GAG GAT TAT GGG TCC TGC AAT TCT ACA GTT TCT TAC TGT TTT GTA 1591/531 1561/521 TCA AAA TCA CTA TCT TTC TGA TAA CAG AAT TGC CAA GGC AGC GGG ATC TCG TAT CTT TAA 1621/541 1651/551 ANA GCA GTC CTC TTA TTC CTA AGG TAA TCC TAT TAA AAC ACA GCT TTA CAA CTT CCA TAT 1681/561 TAC AAA AAA AAA AAA AAA

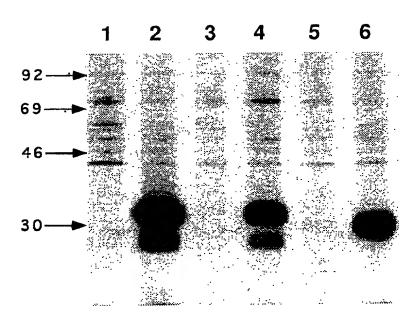


Figure 5

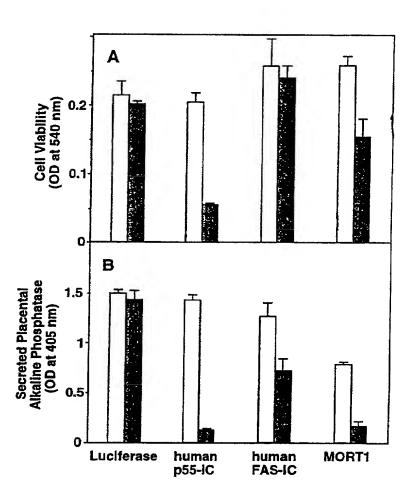


Figure 6

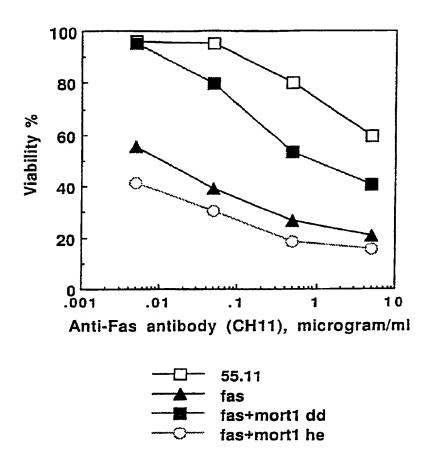


Figure 7